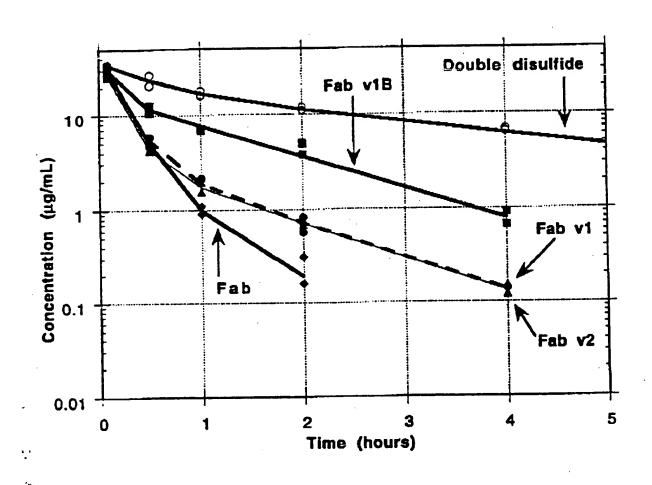
FIGURE 1 A



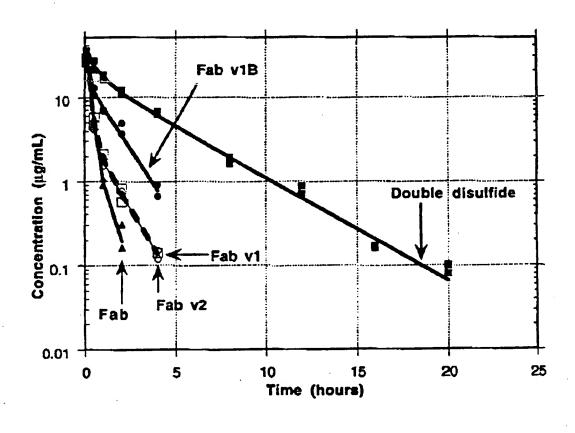


FIGURE 2

	hIgG1	Sequence of human IgG1 CH1 domain	
	hIgG2	Sequence of human IgG2 CH1 domain	
5	hIgG3	Sequence of human IgG3 CH1 domain	
	hIgG4	Sequence of human IgG4 CH1 domain	
	humK	Sequence of human kappa CL domain	
	humL	Sequence of human lambda CL domain	
	=======		
10		114 128 1	3 9
		↓ ↓	\downarrow
	hIgG1	ASTKGPSVFPLAPSSKSTSGGTAA	L
	hIgG2	ASTKGPSVFPLAPCSRSTSESTÄA	L
	hIgG3	ASTKGPSVFPLAPCSRSTSGGTAA	L
15	hIgG4	ASTKGPSVFPLAPCSRSTSESTAA	L
13			
Cold Control and Australia		108 122 13	31
1,:1 1':1		1 1	
## ##	humK	RTVAAPSVFIFPPSDEQLKSGTA	s V
20	humL	Q P K A A P S V T L F P P S S E E L Q A N K A '	ГL
10			
11 to 1	FabCABv1	b ASTKGPSVFPLAPSPKNS SMISNTPA	. L
	of intere	est PKNS SMISNTP	
11		<u></u>	
25	most impo	ortant *** *	
1000 M			
1,3			
	hIgG1	GCLVKDYFPEPVTVSWNSGALTS	
	hIgG2	GCLVKDYFPEPVTVSWNSGALTS	
30	hIgG3	GCLVKDYFPEPVTVSWNSGALTS	
	hIgG4	GCLVKDYFPEPVTVSWNSGALTS	
	humK	V C L L N N F Y P R E A K V Q W K V D N A L Q	
	humL	V C L I S D F Y P G A V T V A W K A D S S P V	
35			
	FabCABv1	b GCLVKDYFPEPVTVSWNSGALTS	
	hIgG1	GVHTFPAVLQSSGLYSLSSV	
	hIgG2	GVHTFPAVLQSSGLYSLSSV	

	hIgG3	GVHTFPAVLQSSGLYSLSSV			
5	hIgG4	$ \texttt{G} \ \texttt{V} \ \texttt{H} \ \texttt{T} \ \texttt{F} \ \texttt{P} \ \texttt{A} \ \texttt{V} \ \texttt{L} \ \texttt{Q} \ \texttt{S} \ \texttt{S} \ \texttt{G} \ \texttt{-} \ \texttt{-} \ \texttt{-} \ \texttt{L} \ \texttt{Y} \ \texttt{S} \ \texttt{L} \ \texttt{S} \ \texttt{S} \ \texttt{V} $			
	humK	SGNSQESVTEQDSKDSTYSLSST			
,	humL	KAGVETTTPSKQSNN-KYAASSY			
			-		
	FabCABv1b	GVHTFPAVLQSSGLYSLSSV			
10					
	•				
	193 200 203				
		\downarrow \downarrow			
	hIgG1	V T V P S S S L G T - Q T Y I C N V N H K P S			
15	hIgG2	V T V P S S N F G T - Q T Y T C N V D H K P S			
12	hIgG3	V T V P S S S L G T - Q T Y T C N V N H K P S			
	hIgG4	V T V P S S S L G T - K T Y T C N V D H K P S			
4 .5		181 190			
		↓			
20	humK	LTLSKADYEKHKVYACEVTHQGL			
(A	humL	LSLTPEQWKSHRSYSCQVTHEGS			
<u> </u>		VI3542G& 2	-		
	FabCABv1b	VTVPHQSLGT-QTYICNVNHKPS			
1 .4	of interes	32 N. 7			
25		- March			
	most impo	rtant			
Ü					
•		·			
30	hIgG1	NTKVDKRV			
	hIgG2	NTKVDKTV			
	hIgG3	N T K V D K R V			
	hIgG4	NTKVDKRV			
35	humK	SSPVTKSFNRGEC			
	humL	TVEKTVAPTECS			
	FabCABv1b	NTKVDKRV			